



#10

R0133045.txt
SEQUENCE LISTING

<110> Kane, Michael D.
Nagel, Aaron C.
Dombkowski, Alan A.

<120> COMPOSITIONS AND SYSTEMS FOR IDENTIFYING AND COMPARING EXPRESSED GENES
(mRNAs) IN EUKARYOTIC ORGANISMS

<130> 65446-87

<160> 5

<170> PatentIn version 3.1

<210> 1

<211> 61

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<220>

<221> misc_feature

<222> (1)..(50)

<223> The poly-T tail may have between 8 and 50 residues.

<220>

<221> misc_feature

<222> (52)..(62)

<223> The sequence may have between 3 and 10 residues selected from a
or c or g or t.

<400> 1
tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt vnnnnnnnnn 60
n 61

<210> 2

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<220>

<221> misc_feature

<222> (23)..(25)

<223> n may be selected from a or c or g or t.

<400> 2
tttttttttt tttttttttt tvnnn 25

<210> 3

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<400> 3
tttttttttt tttttttttt aaac 24

<210> 4

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<400> 4
gtttaaaaaa aaaaaaaaaa aaaa 24

<210> 5

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> This is a synthesized sequence.

<400> 5
tttttttttt tttttttttt ggttt 25